



OIPE

## ENTERED

LUICAEL

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/771,961

DATE: 04/16/2002 TIME: 13:58:26

Input Set : A:\Seqlist.txt

Output Set: N:\CRF3\04162002\1771961.raw

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4 <110> APPLICANT: Donoho, Gregory
      5
              Hilbun, Erin
              Turner, C. Alexander Jr.
      6
              Friedrich, Glenn
      7
              Abuin, Alejandro
      8
     9
              Zambrowicz, Brian
     10
              Sands, Arthur T.
     12 <120> TITLE OF INVENTION: Novel Human BCL-X-Like Proteins and
              Polynucleotides Encoding the Same
     16 <130> FILE REFERENCE: LEX-0127-USA
C--> 18 <140> CURRENT APPLICATION NUMBER: US/09/771,961
C--> 18 <141> CURRENT FILING DATE: 2001-01-29
     18 <150> PRIOR APPLICATION NUMBER: US 60/180,412
     19 <151> PRIOR FILING DATE: 2000-02-04
     21 <160> NUMBER OF SEQ ID NOS: 5
     23 <170> SOFTWARE: FastSEQ for Windows Version 4.0
     25 <210> SEQ ID NO: 1
     26 <211> LENGTH: 984
     27 <212> TYPE: DNA
     28 <213> ORGANISM: Homo sapiens
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     33 gctctcttct caccaaagct gctgagaaca agaagtttgt cccagagggg cctggggaat
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     34 tgttcagcaa atgagtcatg gacagaggtg tcatggcctt gcagaaattc ccaatccagt
     35 gagaaggcca taaaccttgg caagaaaaag tcttcttgga aagcattctt tggagtagtg
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     36 gagaaggaag attcgcagag cacgcctgcc aaggtctctg ctcagggtca aaggacgttg
     37 gaataccaag attegeacag ecageagtgg tecaggtgte titetaaegt ggageagtge
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     38 ttggagcatg aagctgtgga ccccaaagtc atttccattg ccaaccgagt agctgaaatt
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     39 gtttatteet ggeeaceace acaagegaee caggeaggag getteaagte caaagagatt
     40 tttgtaactg agggtctctc cttccagctc caaggccacg tgcctgtagc ttcaagttct
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     41 aagaaagatg aagaagaaca aatactagcc aaaattgttg agctgctgaa atattcagga
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     42 gatcagttgg aaagaaagct gaagaaagat aaggctttga tgggccactt ccaggatggg
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     43 ctgtcctact ctgttttcaa gaccatcaca gaccaggtcc taatgggtgt ggaccccagg
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     44 ggagaatcag aggtcaaagc tcagggcttt aaggctgccc ttgtaataga cgtcacggcc
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     45 aageteacag etattgacaa ecaceegatg aacagggtee tgggetttgg eaccaagtae
     46 ctgaaagaga acttctcgcc atggatccag cagcacggtg gatgggaaaa aatacttggg
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     49 <210> SEQ ID NO: 2
     50 <211> LENGTH: 327
     51 <212> TYPE: PRT
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54 <400> SEQUENCE: 2

52 <213> ORGANISM: Homo sapiens

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55 Met Cys Ser Thr Ser Gly Cys Asp Leu Glu Glu Ile Pro Leu Asp Asp
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57 Asp Asp Leu Asn Thr Ile Glu Phe Lys Ile Leu Ala Tyr Tyr Thr Arg
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59 His His Val Phe Lys Ser Thr Pro Ala Leu Phe Ser Pro Lys Leu Leu
61 Arg Thr Arg Ser Leu Ser Gln Arg Gly Leu Gly Asn Cys Ser Ala Asn
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63 Glu Ser Trp Thr Glu Val Ser Trp Pro Cys Arg Asn Ser Gln Ser Ser
                       70
65 Glu Lys Ala Ile Asn Leu Gly Lys Lys Lys Ser Ser Trp Lys Ala Phe
                                       90
                   85
67 Phe Gly Val Val Glu Lys Glu Asp Ser Gln Ser Thr Pro Ala Lys Val
                                    105
               100
69 Ser Ala Gln Gly Gln Arg Thr Leu Glu Tyr Gln Asp Ser His Ser Gln
                                                    125
                                120
           115
71 Gln Trp Ser Arg Cys Leu Ser Asn Val Glu Gln Cys Leu Glu His Glu
70
                                                140
                            135
       130
73 Ala Val Asp Pro Lys Val Ile Ser Ile Ala Asn Arg Val Ala Glu Ile
                                            155
                        150
75 Val Tyr Ser Trp Pro Pro Pro Gln Ala Thr Gln Ala Gly Gly Phe Lys
                                        170
                    165
 77 Ser Lys Glu Ile Phe Val Thr Glu Gly Leu Ser Phe Gln Leu Gln Gly
                                                        190
                                    185
                180
 79 His Val Pro Val Ala Ser Ser Lys Lys Asp Glu Glu Glu Gln Ile
 81 Leu Ala Lys Ile Val Glu Leu Leu Lys Tyr Ser Gly Asp Gln Leu Glu
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                            215
 83 Arg Lys Leu Lys Lys Asp Lys Ala Leu Met Gly His Phe Gln Asp Gly
                        230
 85 Leu Ser Tyr Ser Val Phe Lys Thr Ile Thr Asp Gln Val Leu Met Gly
                                         250
                    245
 87 Val Asp Pro Arg Gly Glu Ser Glu Val Lys Ala Gln Gly Phe Lys Ala
                                     265
                260
 89 Ala Leu Val Ile Asp Val Thr Ala Lys Leu Thr Ala Ile Asp Asn His
                                 280
 91 Pro Met Asn Arg Val Leu Gly Phe Gly Thr Lys Tyr Leu Lys Glu Asn
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 93 Phe Ser Pro Trp Ile Gln Gln His Gly Gly Trp Glu Lys Ile Leu Gly
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 100 <212> TYPE: DNA
 101 <213> ORGANISM: Homo sapiens
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 105 accatagaat tcaaaatcct cgcctactac accagacatc atgtcttcaa gagcacccct
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106 gctctctct caccaaagct gctgagaaca agaagtttgt cccagagggg cctggggaat 107 tgttcagcaa atgagtcatg gacagaggtg tcatggcctt gcagaaattc gcaatccagt 108 gagaaggcca taaaccttgg caagaaaaag tcttcttgga aagcattctt tggagtagtg 109 gagaaggaag attcgcacag cacgcctgce caagcagtgg tccaggtgt tctctaggt aaggacgttg 110 gaataccaag attcgcacag cccaaaagtc attccattg ggagacagtgt tccaggtgtc ttctaacgt ggagcagtgc 111 ttggagcatg aagctgtgga ccccaaagtc attccattg ccaaccgagt agctgaaatt 112 gtttattcct ggccaccacc acaagcgacc cttccaggtc caaggcaggag gcttcaagtc caaaggaatt ttgtaactg agggtctctc cttccagctc caaggccacg acaaccgagt tccaaggtc tccaaggtc ccaaaccgagt tccaaggtc ccaaaccgagt agctgaaatt caaaggaatt tccattg ccaaccgagt agctgaaatt aggacaccg acaaggcaccg aaaaattgttg agctgctgaa atattcagga cactgccttc ttgatggc tttgatggc ccttggttga caccagcatc ttgattga caccagcatc tgcctttgat caccagcatc tgcctttgat caccagcatc tgcatttga caccagcatc tgcatttga caccagcatc ttgattga caccagcatc tgcatttga caccagcacc aaccagcatc tgcatttga caccagcaccaccaccaccaccaccaccaccaccaccacc	180 240 300 360 420 480 540 600 660 720 759
124 Met Cys Ser Thr Ser Gly Cys Asp Leu Glu Glu He Flo Heu Asp Hop	
126 Asp Asp Leu Asn Thr Ile Glu Phe Lys IIe Leu Ala 191 191 111 1119	
127 20 25 128 His His Val Phe Lys Ser Thr Pro Ala Leu Phe Ser Pro Lys Leu Leu	
129 35 40 130 Arg Thr Arg Ser Leu Ser Gln Arg Gly Leu Gly Asn Cys Ser Ala Asn	
131 50 53 132 Glu Ser Trp Thr Glu Val Ser Trp Pro Cys Arg Asn Ser Gln Ser Ser	
70 13	
134 Glu Lys Ala Ile Asn Leu Gly Lys Lys Lys Ser Ser IIp Lys Ala Inc	
135 136 Phe Gly Val Val Glu Lys Glu Asp Ser Gln Ser Thr Pro Ala Lys Val	
137 100 103 138 Ser Ala Gln Gly Gln Arg Thr Leu Glu Tyr Gln Asp Ser His Ser Gln	
446 170	٠
139 115 120 120 120 140 Gln Trp Ser Arg Cys Leu Ser Asn Val Glu Gln Cys Leu Glu His Glu	
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141 130 133 133 133 142 Ala Val Asp Pro Lys Val Ile Ser Ile Ala Asn Arg Val Ala Glu Ile 142 Ala Val Asp Pro Lys Val Ile 155 160	
143 145 144 Val Tyr Ser Trp Pro Pro Pro Gln Ala Thr Gln Ala Gly Gly Phe Lys	
145 146 Ser Lys Glu Ile Phe Val Thr Glu Gly Leu Ser Phe Gln Leu Gln Gly	
100 187	
147 180 105 105 148 His Val Pro Val Ala Ser Ser Ser Lys Lys Asp Glu Glu Glu Gln Ile	
149 195 200 150 Leu Ala Lys Ile Val Glu Leu Leu Lys Tyr Ser Gly Asp Gln Leu Glu 215 220	
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152 225	
154 Gln Gly Phe Pro Gln Asp Gly Leu Met Ala Cys 11e	
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157 <210> SEQ ID NO: 5	

RAW SEQUENCE LISTING

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Input Set : A:\Seqlist.txt

Output Set: N:\CRF3\04162002\1771961.raw

158 <211> LENGTH: 2132		
159 <212> TYPE: DNA		
160 <213> ORGANISM: Homo sapiens		
162 <400> SEQUENCE: 5		
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165 atagaaatat cottactgoo acctgacctg	aaqcaqaaqa aatcacagac agcttccaga	180
166 ccaggcccaa catgtgtagc accagtgggt	gtgacctgga agaaatcccc clayalyaly	240
167 atgacctaga caccatagaa ttcaaaatcc	togoctacta caccagacat calgicilea	300
168 agaggagged toctetete teaccaaage	tgctgagaac aagaagtttg tcccagaggg	360
160 gootgaggaa ttgttcagga aatgagtcat	ggacagaggt gtcatggcct tgcagaaatt	420
170 occaatorag tgagaaggcc ataaaccttg	qcaaqaaaaa gtcttcttgg aaagcattct	480
171 ++ggagtagt ggagaaggaa gattcgcaga	gcacgcctgc caaggtctct gctcagggtc	540
172 agaggacgtt ggaataccaa gattcgcaca	qccaqcagtg gtccaggtgt ctttctaacg	600
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176 ottoaagtto taagaaagat gaagaagaac	aaatactagc caaaattgtt gagcigciga	840
177 aatattcagg agatcagttg gaaagaaagg	acactgoott catocccatt coottggily	900
178 acaccagcat ccagggtttt ccacaggatg	gtttgatggc ctgcatttga gctaaagaac	960
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183 acaaccaccc gatgaacagg gtcctgggct	ttggcaccaa gtacctgaaa gagaacttct	1260
184 caccatagat ccagcagcac ggtggatggg	aaaaaatact tgggatatca calyaayaay	1320
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187 ttcaaaacca ttattcctgt gactggagag	qcatcaggag agggctcgtt cgtctccagc	1560
188 teatagaatg tagcagcate atcettgaca	gtgatgtttt tcaggccctc cattgagaac	1620
189 ctgaggaaat ctgtaaagat aagtggtgat	gttqtttcaa acgttcagaa cagalaccal	1680
190 catcctgcct ttgttagctg ctgtagggaa	agtgcgttac agatgtctgc tgacctcaca	1740
191 agagtgaaaa gataaactgt gcatgtgttt	ccaaaaacgt ttctagtact attlaction	1800
100 asactacact togggtggcc taatacctag	gaagatgttg ctattcacgt tagtaaacag	1860
193 cctaaagaaa ctcttaggtt tactgctaca	tccatttgtt tggagaggta actgttgtct	1920
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195 ttttctcagg gaccacacct cttcttccca	aggtccctgg gacttcctca ttctttgtgg	2040
196 tagtacaatg attggtagca ggtaaaataa	atacatagaa agacwamwrw caaaagagig	2100
197 tcttctgatt agtaaggaat aagtcttctg	gattatcaaa tggagttaat ttacataaaa	2100
198 atgctcaaaa cacttcttgg tacttaatgt	; ta	4134

VERIFICATION SUMMARY

DATE: 04/16/2002

PATENT APPLICATION: US/09/771,961

TIME: 13:58:28

Input Set : A:\Seqlist.txt

Output Set: N:\CRF3\04162002\I771961.raw

L:18 M:270 C: Current Application Number differs, Replaced Current Application No L:18 M:271 C: Current Filing Date differs, Replaced Current Filing Date